

LOCUS AK001486 2954 bp mRNA PRI 22-FEB-2000
 DEFINITION Homo sapiens cDNA FLJ10624 fis, clone NT2RP2005525, highly similar to Mus musculus kanadaplin mRNA.
 ACCESSION AK001486
 VERSION AK001486.1 GI:7022771
 KEYWORDS oligo capping; fis (full insert sequence).
 SOURCE Homo sapiens teratocarcinoma cell line:NT2 cDNA to mRNA, clone_lib:NT2RP2 clone:NT2RP2005525.
 ORGANISM Homo sapiens
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
 REFERENCE 1 (sites)
 AUTHORS Isogai,T., Ota,T., Hayashi,K., Sugiyama,T., Otsuki,T., Suzuki,Y., Nishikawa,T., Nagai,K., Sugano,S., Ishibashi,T., Fujimori,K., Tanai,H., Kimata,M., Watanabe,M., Hiraoka,S., Ishii,S., Kawai,Y., Saito,K., Yamamoto,J., Wakamatsu,A., Nakamura,Y., Nagahari,K., Masuho,Y. and Kanehori,K.
 TITLE NEDO human cDNA sequencing project
 JOURNAL Unpublished (2000)
 REFERENCE 2 (bases 1 to 2954)
 AUTHORS Isogai,T. and Otsuki,T.
 TITLE Direct Submission
 JOURNAL Submitted (16-FEB-2000) to the DDBJ/EMBL/GenBank databases. Takao Isogai, Helix Research Institute, Genomics Laboratory; 1532-3 Yana, Kisarazu, Chiba 292-0812, Japan (E-mail:genomics@hri.co.jp, Tel:81-438-52-3951, Fax:81-438-52-3952)
 COMMENT NEDO human cDNA sequencing project supported by Ministry of International Trade and Industry of Japan; cDNA full insert sequencing: Research Association for Biotechnology; cDNA library construction, 5'- & 3'-end one pass sequencing and clone selection: Helix Research Institute (supported by Japan Key Technology Center etc.) and Department of Virology, Institute of Medical Science, University of Tokyo.
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BASE COUNT 894 a 639 c 776 g 645 t
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LOCUS BAA91718 796 aa PRI 22-FEB-2000
 DEFINITION unnamed protein product [Homo sapiens].
 ACCESSION BAA91718
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 ORGANISM Homo sapiens
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
 Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
 REFERENCE 1 (sites)
 AUTHORS Isogai,T., Ota,T., Hayashi,K., Sugiyama,T., Otsuki,T., Suzuki,Y.,
 Nishikawa,T., Nagai,K., Sugano,S., Ishibashi,T., Fujimori,K.,
 Tanai,H., Kimata,M., Watanabe,M., Hiraoka,S., Ishii,S., Kawai,Y.,
 Saito,K., Yamamoto,J., Wakamatsu,A., Nakamura,Y., Nagahari,K.,
 Masuho,Y. and Kanehori,K.
 TITLE NEDO human cDNA sequencing project
 JOURNAL Unpublished (2000)
 REFERENCE 2 (residues 1 to 796)
 AUTHORS Isogai,T. and Otsuki,T.
 TITLE Direct Submission
 JOURNAL Submitted (16-FEB-2000) to the DDBJ/EMBL/GenBank databases. Takao
 Isogai, Helix Research Institute, Genomics Laboratory; 1532-3 Yana,
 Kisarazu, Chiba 292-0812, Japan (E-mail:genomics@hri.co.jp,
 Tel:81-438-52-3951, Fax:81-438-52-3952)
 COMMENT NEDO human cDNA sequencing project supported by Ministry of
 International Trade and Industry of Japan; cDNA full insert
 sequencing; Research Association for Biotechnology; cDNA library
 construction, 5'- & 3'-end one pass sequencing and clone selection:
 Helix Research Institute (supported by Japan Key Technology Center
 etc.) and Department of Virology, Institute of Medical Science,
 University of Tokyo.
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